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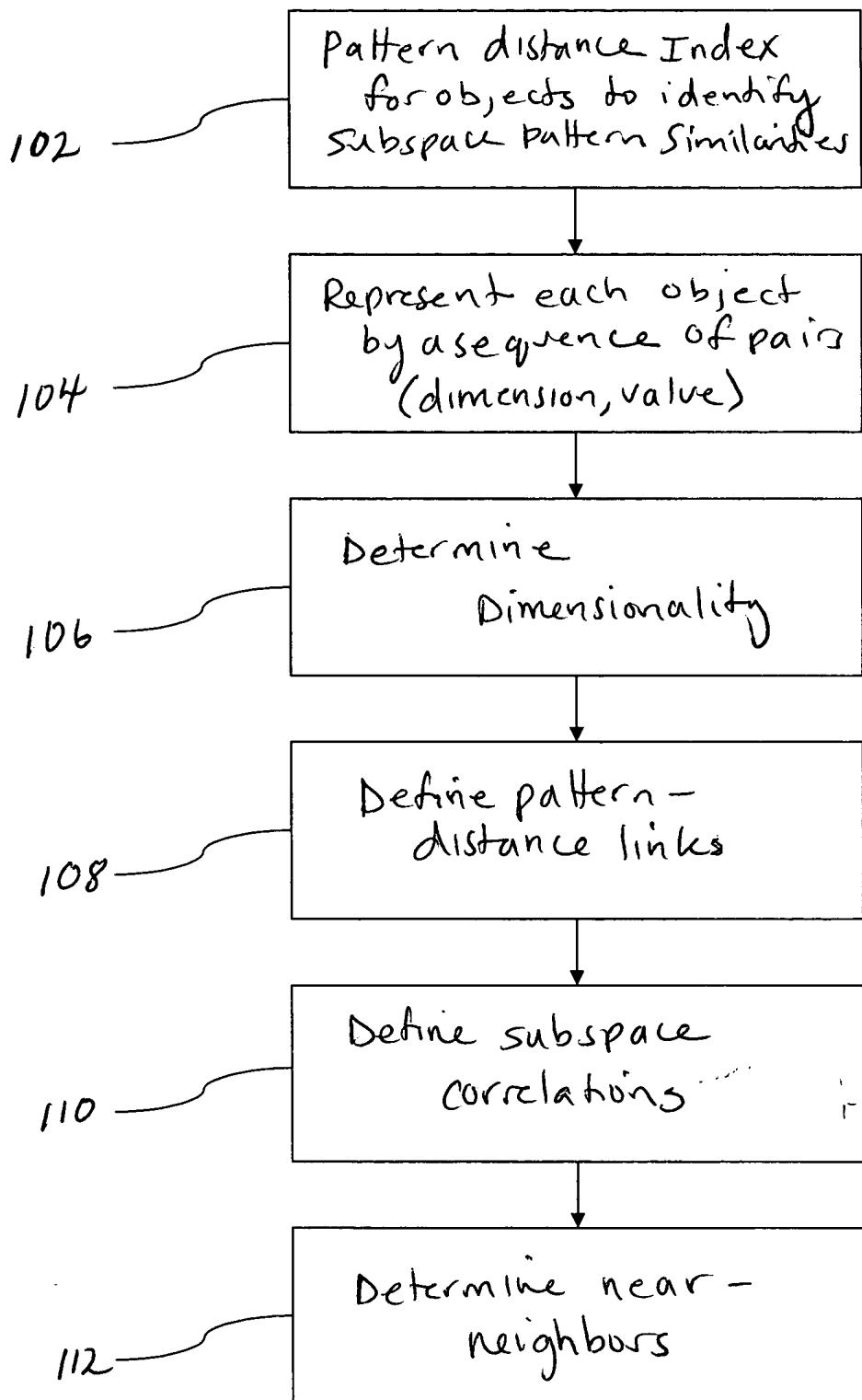


FIG. 1

200

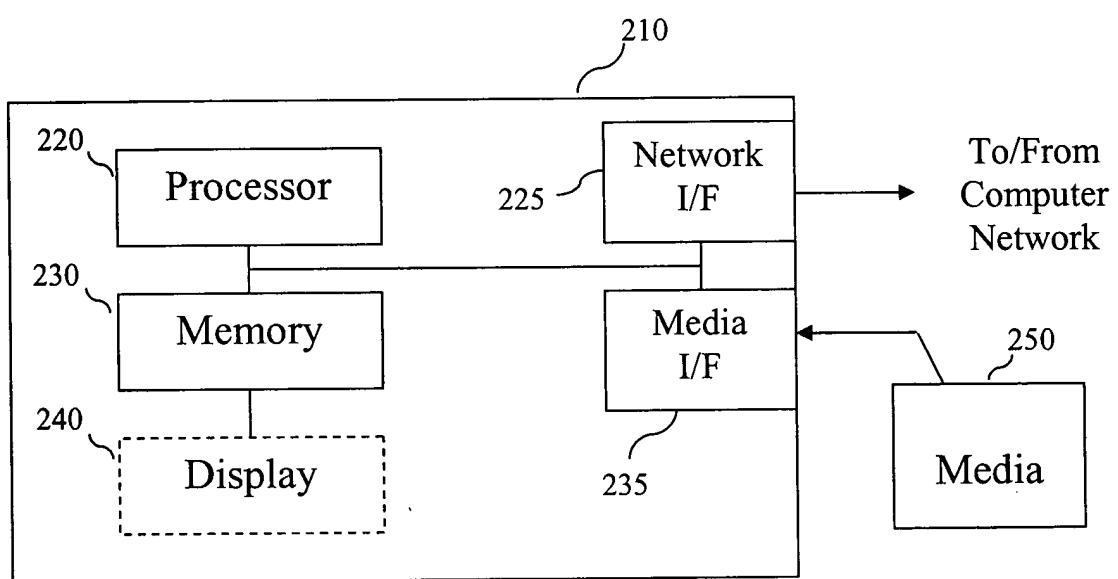


FIG. 2

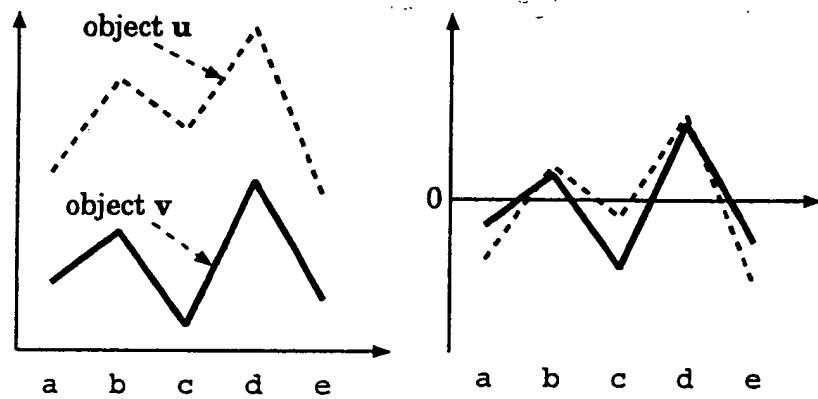


FIG. 3

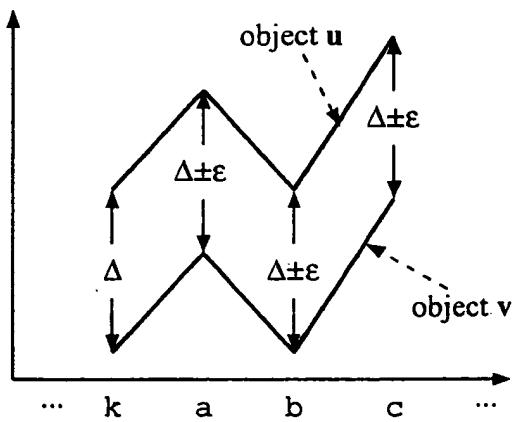


FIG. 4

$f(u, i)$, where $u \in \{\#1, \#2\}$ and $i = 1, \dots, 4$				
$(c_1, 0)$,	$(c_2, -3)$,	$(c_3, 1)$,	$(c_4, -1)$,	$(c_5, -3)$
$(c_2, 0)$,	$(c_3, 4)$,	$(c_4, 2)$,	$(c_5, 0)$	
$(c_3, 0)$,	$(c_4, -2)$,	$(c_5, -4)$		
	$(c_4, 0)$,	$(c_5, -2)$		
$(c_1, 0)$,	$(c_2, -3)$,	$(c_3, 1)$,	$(c_4, -1)$,	$(c_5, 2)$
$(c_2, 0)$,	$(c_3, 4)$,	$(c_4, 2)$,	$(c_5, 5)$	
$(c_3, 0)$,	$(c_4, -2)$,	$(c_5, 1)$		
	$(c_4, 0)$,	$(c_5, 3)$		

FIG. 5

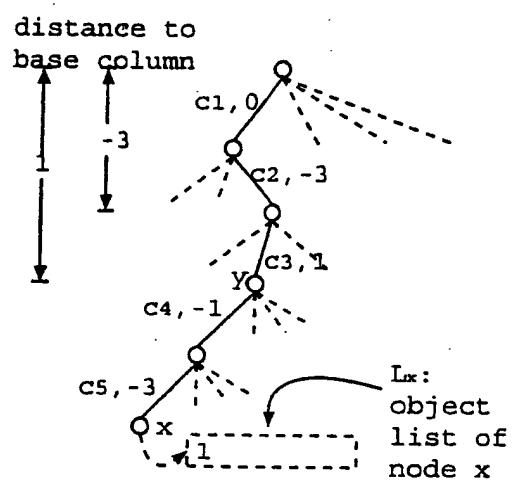


FIG. 6

Input: T : a trie built on \mathcal{D}
 S : a subspace defined by a continuous column
set $\{c_i, c_{i+1}, \dots, c_k\}$
 $q = (c_1, v_1), \dots, (c_n, v_n)$: a query object
 ϵ : pattern threshold

Output: near-neighbors of q in subspace S

$n \leftarrow$ root of T ;
 $search(n, S)$;

Function $search(x, S)$

if $S = \emptyset$ **then**
| output the descendants of x ;

else
| assume $S = \{c_j, c_{j+1}, \dots, c_k\}$;
| **for** x 's child node y under edge labeled (c_j, v)
| where $v \in [(v_j - v_i) - \epsilon, (v_j - v_i) + \epsilon]$ **do**
| | $search(y, \{c_{j+1}, \dots, c_k\})$;

FIG. 7

Input: \mathcal{D} : objects in multi-dimensional space \mathcal{A}

Output: PD-Index of \mathcal{D}

for each $u \in \mathcal{D}$ **do**

\lfloor insert $f(u, i)$, $1 \leq i < |\mathcal{A}|$ into a trie; (Eq 5)

for each node x encountered in a depth-first traversal of
the trie **do**

\lfloor label node x by $\langle n_x, s_x \rangle$;

\lfloor let (c, d) be the arc that points to x ;

\lfloor append $\langle n_x, s_x \rangle$ to pattern-distance link (c, d) ;

FIG 8

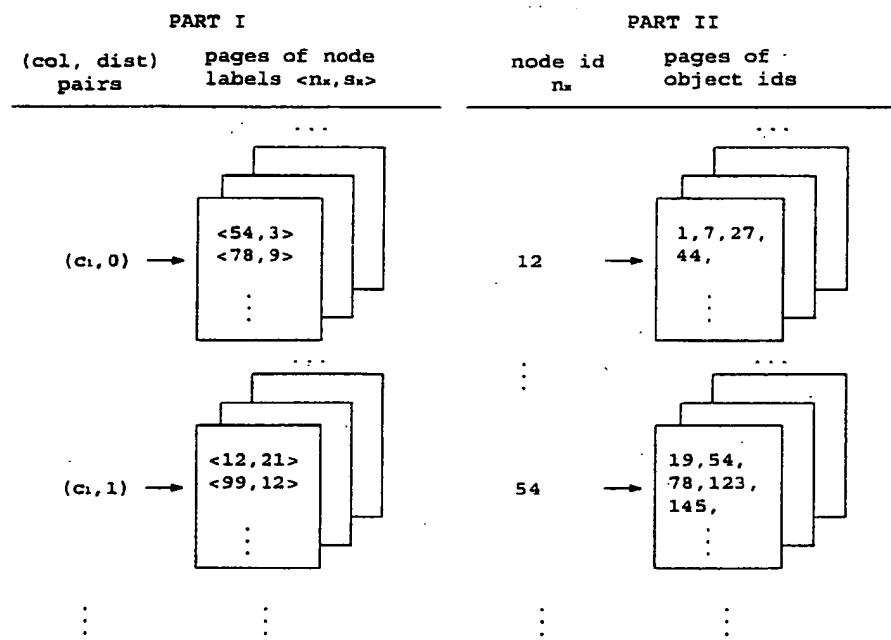


FIG. 9A

FIG. 9B

Input: q : a query object, S : a given subspace
 ϵ : pattern threshold

Output: q 's near-neighbors in subspace S

let $(c_1, v_1), \dots, (c_{|S|}, v_{|S|})$ be q 's projection on S ;
 $x \leftarrow$ the node under arc $(c_1, 0)$;
 $search(x, 2)$;

Function $search(x, i)$

if $i \leq |S|$ **then**

for pattern distance link I of (c_i, v) , where $v \in [v_i - v_1 - \epsilon, v_i - v_1 + \epsilon]$ **do**

 /* perform a binary search on I */

for all node $r \in I$ and $n_r \in [n_x, n_x + s_x]$ **do**

 | $search(r, i + 1)$;

 | **end**

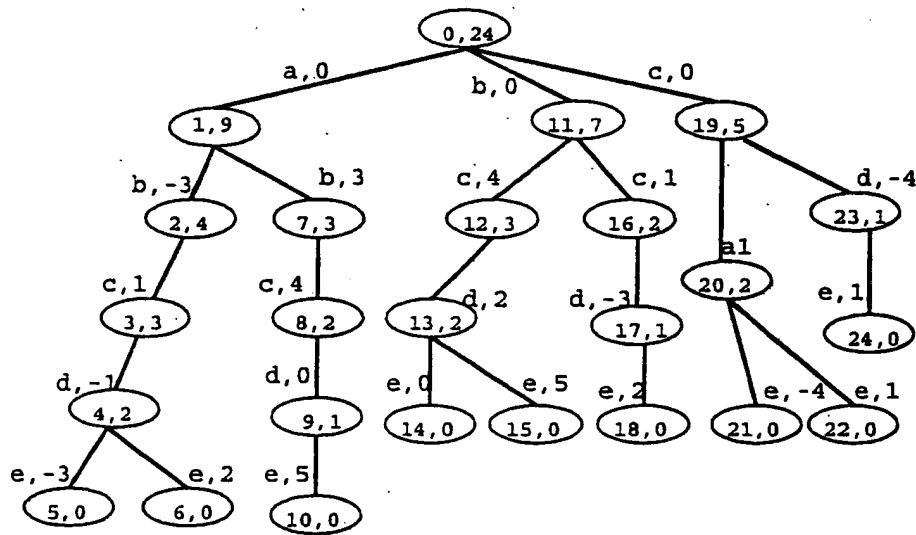
 | **end**

 | **else**

 | output objects in L_x , $x = v_s, \dots, v_m$

 | **end**

FIG. 10



node	5	6	10	14	15	18	21	22	24
objs	{1}	{2}	{3,4}	{1}	{2}	{3,4}	{1}	{2}	{3,4}

FIG. 11

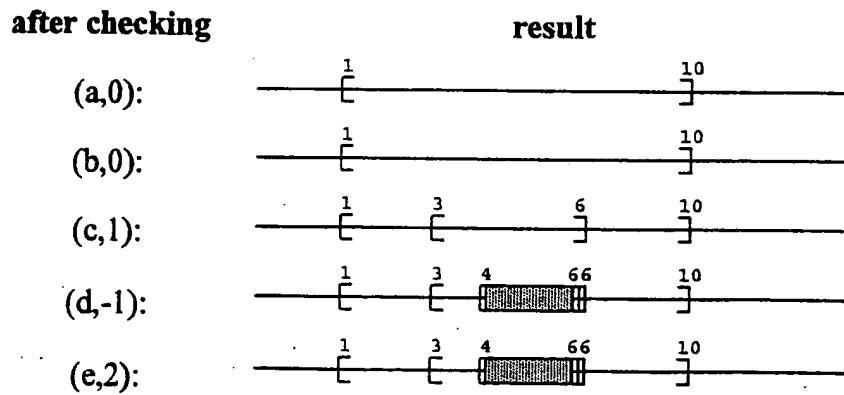


FIG. 12

Input: $q = (c_1, v_1), \dots, (c_n, v_n)$: a query object
 r : distance threshold, ϵ : pattern tolerance
 F : index file for \mathcal{D}

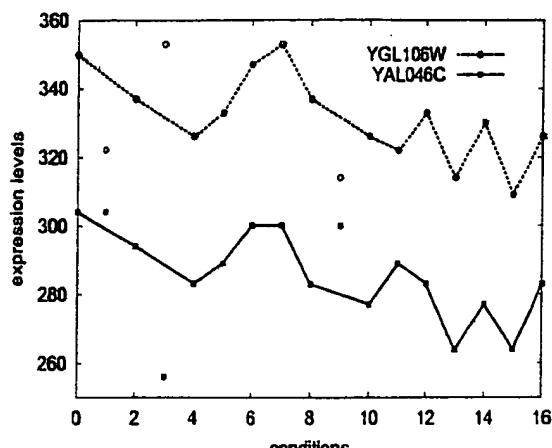
Output: $\mathcal{NN}(q, r)$

```

for  $i = 1, \dots, r + 1$  do
   $R \leftarrow$  the range of the (only) node in link  $(c_i, 0)$ ;
   $j \leftarrow i + 1$ ;
  while  $R \neq \phi$  and  $j \leq |\mathcal{A}|$  do
    search link  $(c_j, v)$  for nodes inside any range of
     $R$ , where  $v \in [v_j - v_i - \epsilon, v_j - v_i + \epsilon]$ ;
    update  $R$  by adding the ranges of those nodes;
    if a region  $s$  of  $R$  is inside  $|\mathcal{A}| - r$  brackets then
      output objects in  $L_x$  where  $x \in s$ ;
      eliminate  $s$  from  $R$ ;
    end
    if a region  $s$  of  $R$  is inside less than  $r - j$  brackets then
      eliminate the region from  $s$ ;
    end
     $j \leftarrow j + 1$ ;
  end
end

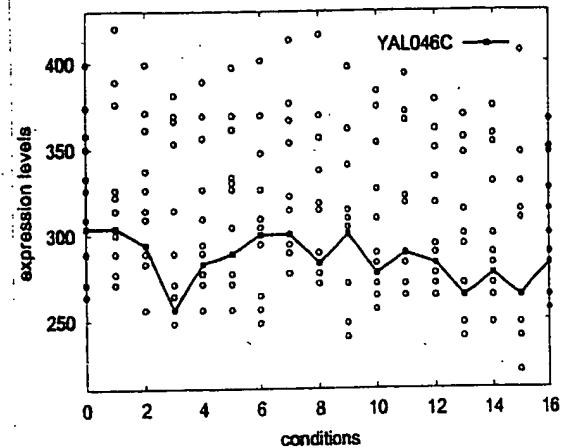
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FIG. 13



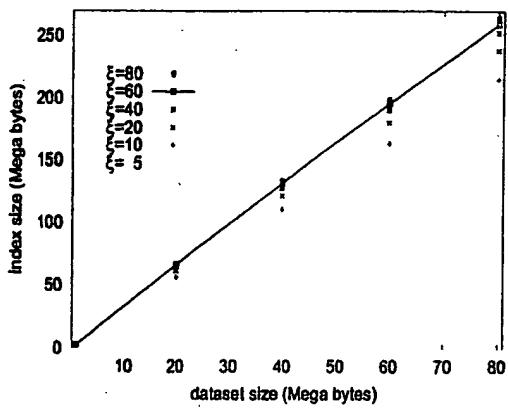
$pdist(\cdot, \cdot) \leq 3$

FIG.14A

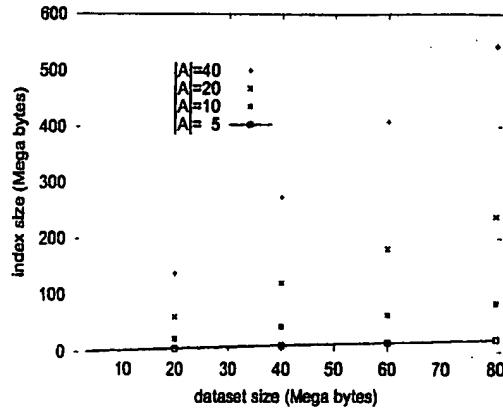


$pdist(\cdot, \cdot) \leq 4$

FIG.14B



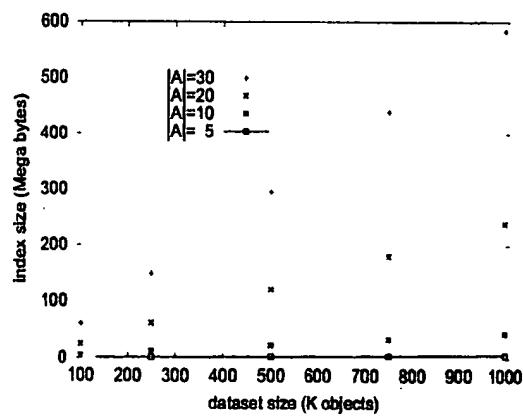
$|A| = 20, \xi = 5, \dots, 80$



varying total data size, $\xi = 20$

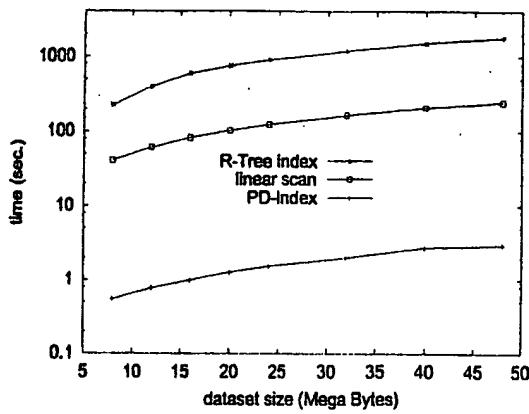
FIG. 15A

FIG. 15B

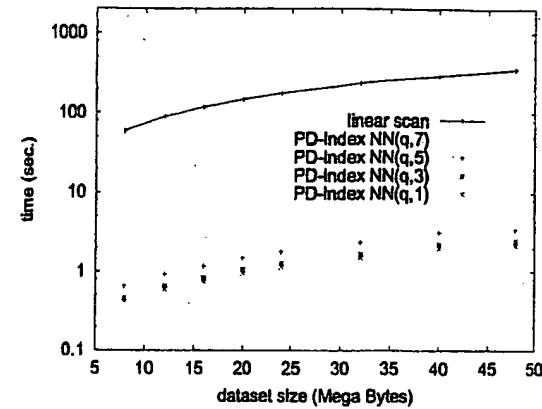


varying # of objects, $\xi = 20$

FIG. 15c



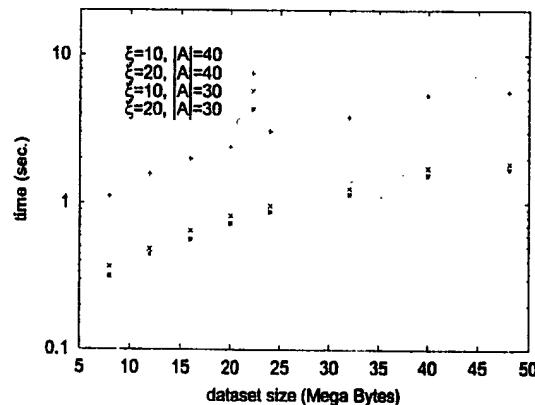
Pattern matching in given subspaces



Near-neighbor search in subspaces beyond given dimensionalities

FIG. 16A

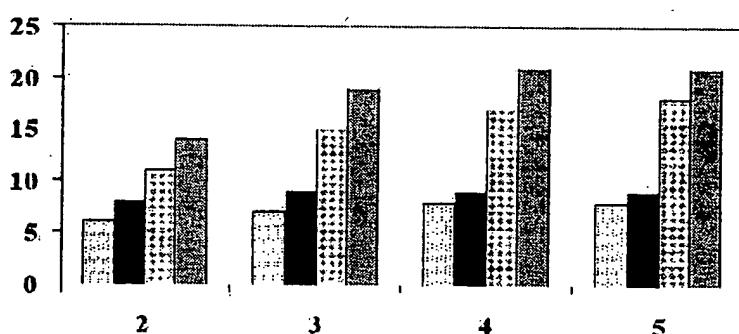
FIG. 16B



Impact of ξ and $|\mathcal{A}|$ in near-neighbor query $\mathcal{NN}(q, 7)$

FIG. 16C

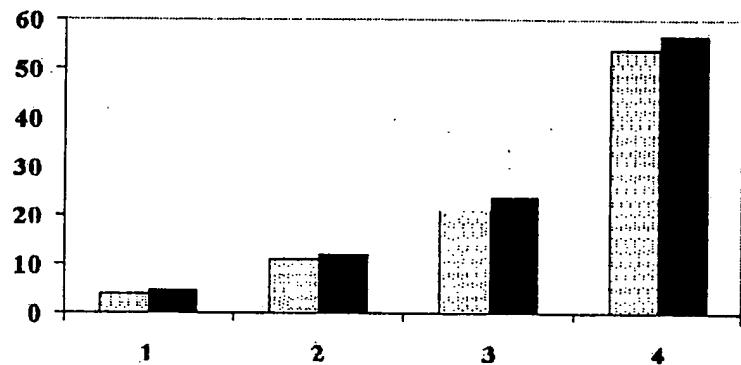
■ # Nodes Yeast ■ # Pages Yeast.
■ # Nodes Mouse ■ # Pages Mouse



(a) Find Near-neighbors in DNA micro-array
in given subspaces (X axis is query length)

FIG. 17A

■ # Nodes ■ # Pages



(b) Find Near-Neighbors in DNA micro-array
(X axis is the distance radius r)

FIG. 17B